

Claims

1. A method for processing information on a nucleotide sequence comprising steps of:

(a) obtaining positional information representing a position in a nucleotide sequence in accordance with a request for an object and/or service; and

(b) evaluating adequacy of transmission of nucleotide sequence-related information corresponding to the positional information obtained in step (a), based on the flag information associated with the positional information for evaluating adequacy of transmission of nucleotide sequence-related information associated with the positional information representing a position in a nucleotide sequence.

2. The method for processing information on a nucleotide sequence according to claim 1, which further comprises steps of:

(c) obtaining the nucleotide sequence-related information that was evaluated adequate for transmission in step (b) from among nucleotide sequence-related information; and

(d) transmitting the nucleotide sequence-related information obtained in step (c).

3. The method for processing information on a nucleotide sequence according to claim 1, wherein processing is cancelled when some of the positional information obtained in step (a) correspond to the nucleotide sequence-related information that was evaluated inadequate for transmission in step (b).

4. A method for processing information on a nucleotide sequence comprising steps of:

(a) obtaining positional information representing a position in a nucleotide sequence in accordance with a request for an object and/or service;

(b) obtaining flag information associated with the positional information and/or information concerning the flag information, wherein the flag information is for evaluating adequacy of transmission of nucleotide sequence-related information associated with the positional information representing a position in a nucleotide sequence; and

(c) evaluating adequacy of transmission of nucleotide sequence-related information corresponding to the positional information obtained in step (a), based on the flag information and/or information concerning the flag information obtained in step (b).

5. The method for processing information on a nucleotide sequence according to claim 4, which further comprises steps of:

(d) obtaining the nucleotide sequence-related information that was evaluated adequate for transmission in step (c) from among nucleotide sequence-related information; and

(e) transmitting the nucleotide sequence-related information obtained in step (d).

6. The method for processing information on a nucleotide sequence according to claim 4, wherein processing is cancelled when some of the positional information obtained in step (a) correspond to the nucleotide sequence-related information that was evaluated inadequate for transmission in step (c).

7. A method for processing information on a nucleotide sequence comprising steps of:

(a) evaluating adequacy of transmission of nucleotide sequence-related information, based on the flag information associated with positional information for evaluating adequacy of transmission of nucleotide sequence-related information associated with the positional information representing a position in a nucleotide sequence;

(b) obtaining the nucleotide sequence-related information that was evaluated adequate for transmission in step (a) from among nucleotide sequence-related information;

(c) transmitting the nucleotide sequence-related information obtained in step (b);
and

(d) transmitting request information for an object and/or service.

8. A method for processing information on a nucleotide sequence comprising steps of:

(a) obtaining flag information associated with the positional information and/or information concerning the flag information, wherein the flag information is for evaluating adequacy of transmission of nucleotide sequence-related information associated with the positional information representing a position in a nucleotide sequence;

(b) evaluating adequacy of transmission of nucleotide sequence-related information based on the flag information and/or information concerning the flag information obtained in step (a);

(c) obtaining the nucleotide sequence-related information that was evaluated adequate for transmission in step (b) from among nucleotide sequence-related information;

(d) transmitting the nucleotide sequence-related information obtained in step (c);
and

(e) transmitting request information for an object and/or service.